



S2 Fig. Comparison of SPR determined $\Delta\Delta G$ with the deep mutagenesis binding data from Procko and co-workers²². A. Experimental $\Delta\Delta G$ vs. ACE2 mutant enrichment in the high RBD binding population. The regression parameters are: slope = 0.38, intercept = -0.14, $R^2 = 0.59$ and $p = 0.01$. B. Experimental $\Delta\Delta G$ vs. ACE2 mutant enrichment in the low RBD binding population (negated). The regression parameters are: slope = 0.41, intercept = -0.18, $R^2 = 0.74$ and $p = 0.002$. Figure generated with R ggplot2.